## IE 529 HW3 Zhenye Na Zna2

1. Use induction to prove Jensen's Inequality, as follows: \frac{2}{2} \pi\_{\text{x}} \text{x}; \left\{\frac{2}{2} \pi\_{\text{x}}; \text{x};}\right\}

Base case:

For n=1,2, the equality is true.

 $\alpha_1 f(x_1) + \alpha_2 f(x_2) \leq f(\alpha_1 X_1 + \alpha_2 X_2)$ 

Induction steps:

Assume that n=m, the inequality stays true.

Then for n=m+1:

as dent is not related to i, so -> dentif(Xent) + f [ = d; x; ]

> dentif(Xent) + = d; d; f(x;)

= <u>\frac{2}{121}</u> \times; \frac{1}{1}(\times\_i)

Equality holds if and only if  $X_1 = X_2 = \cdots = X_n$ , or f(x) is linear.

2. We have noticed that  $f(x) = \log(x)$  is concave on  $(0, \infty)$  So, we take  $\log$  on the left-hand side of the inequality.  $\log(\frac{\pi}{1!}x_i)^{\frac{1}{n}} = \frac{1}{n!}(\log x_i + \log x_2 + \dots + \log x_n) = \frac{1}{n!} + \log x_i \leq \log(\frac{\pi}{1!} + x_i)$ Be cause  $\begin{cases} x_i \\ \vdots \\ x_{i-1} \end{cases}$  is a non-negative set,  $f(x) = \log(x)$   $\begin{cases} x_i \\ \vdots \\ x_{i-1} \end{cases} \times i \end{cases} \leq (\frac{\pi}{n} + \frac{\pi}{n}) \times i$ 

3.  $Y = \beta x + e$ 

$$Y = \begin{bmatrix} y_1 \\ y_2 \\ y_3 \\ \vdots \\ y_n \end{bmatrix} \qquad X = \begin{bmatrix} X_1 \\ X_2 \\ X_3 \\ \vdots \\ X_n \end{bmatrix} \quad \text{and} \quad e = \begin{bmatrix} e_1 \\ e_2 \\ e_3 \\ \vdots \\ e_n \end{bmatrix}$$

and 
$$\beta = [\beta]$$
,  $Y = X\beta + e$ 

The least-squares line minimizes

$$Q(\beta) = \sum_{i=1}^{n} (y_i - \beta x_i)^2 = (Y - X\beta)^T (Y - X\beta)$$

The least squares estimate  $\beta$  solves the first order equation:  $\frac{\partial Q(\beta)}{\partial \beta} = 0$  and is given by

$$\hat{\beta} = (X^T X)^T X^T Y = \hat{S}_{i=1} (X_i^2)^{-1} \hat{S}_{i=1} \times_i y_i = \frac{\hat{S}_{i=1}}{\hat{S}_{i=1}} \chi_i^2 y_i$$

(b)  $\hat{\beta} = \frac{\hat{S}}{\hat{S}} W_i Y_i$ , where  $W_i = \frac{X_i}{\sum_{j=1}^n x_j^2}$ . It can be considered as a weighted sum of the independent normal random variables :  $y_i \sim N(x_i \beta, \sigma^2)$ , which is a normal distribution. So Next step is to figure out  $M_{\beta}$  and  $\sigma_{\beta}^2$ .

$$E(\hat{\beta}) = E(\hat{\xi}, \omega; y_i)$$

$$= \hat{\xi}_{i=1}^2 \omega_i E(y_i) = \hat{\xi}_{i=1}^2 \omega_i \cdot (x_i \beta)$$

$$= \hat{\xi}_{i=1}^2 \frac{x_i}{\hat{\xi}_{i}} \cdot x_i \beta$$

$$= \beta \cdot \hat{\xi}_{i=1}^2 \frac{x_i}{\hat{\xi}_{i}} \cdot x_i \beta$$

$$= \beta \cdot \hat{\xi}_{i=1}^2 \frac{x_i}{\hat{\xi}_{i}} \cdot x_i \beta$$

$$Var(\hat{\beta}) = Var(\hat{\xi}, \omega; y_i)$$

$$\begin{aligned}
& \bigvee_{\alpha \Gamma} (\hat{\beta}) = \bigvee_{\alpha \Gamma} \left( \frac{\hat{\Sigma}}{i=1} \omega_{i} y_{i} \right) \\
& = \underbrace{\hat{\Sigma}}_{i=1} \omega_{i}^{2} \cdot \bigvee_{\alpha \Gamma} (y_{i}) = \underbrace{\hat{\Sigma}}_{i=1} \omega_{i}^{2} \cdot \sigma^{2} \\
& = D^{2} \times \underbrace{\hat{\Sigma}}_{i=1} \left[ \left( \frac{x_{i}^{2}}{\sum_{j=1}^{n} x_{j}^{2}} \right)^{2} \right] \\
& = D^{2} \times \underbrace{\frac{\sum_{i=1}^{n} x_{i}^{2}}{\left( \sum_{j=1}^{n} x_{j}^{2} \right)^{2}}}_{= D^{2} \times \underbrace{\frac{1}{\sum_{i=1}^{n} x_{i}^{2}}}_{= D^{2} \times \underbrace{\frac{1}{\sum_{i=1$$

 $= \overline{\Sigma}^{2} \times \frac{1}{\overline{\Sigma}_{j}^{2} \cdot 1} \times \frac{1}{\overline$ 

(c). 
$$SS_R = \frac{\hat{S}}{1-1} (y_1 - x_1 \hat{\beta})^2 \sim \sigma^2 \chi_{(n-1)}^2$$

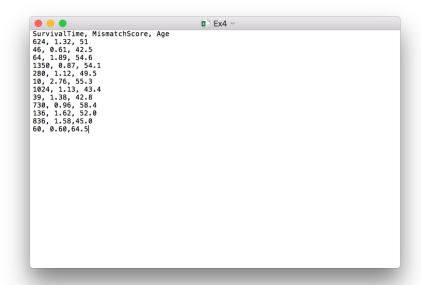
(d)

A significance level 
$$\frac{x}{(n-2)\frac{S_{xx}}{SC_R}}$$
 test of  $H_0$ :  
reject  $H_0$  if  $\frac{(n-2)\frac{S_{xx}}{SC_R}}{SC_R}$   $|B| > t_{\frac{x}{2},n-1}$ 

accept Ho otherwise

rejecting Ho if the desired significance level is at least as large as:

## **Problem 4:**



Below is the R program used for this problem.

```
# read csv file
mydata <- read.csv("/Users/macbookpro/Desktop/Ex4.csv")</pre>
x1<- mydata$MismatchScore;
x2<- mydata$Age;
t <- mydata$SurvivalTime;
y < - log2(t);
#fit log model
fit <-lm(y \sim x1 + x2)
#Results of the model
summary(fit)
             Call:
             lm(formula = y \sim x1 + x2)
             Residuals:
                      1Q Median
                                   3Q
               Min
                                         Max
             -3.481 -1.815 0.166 1.799 2.402
             Coefficients:
                        Estimate Std. Error t value Pr(>|t|)
             (Intercept) 11.5724
                                  5.3701 2.15
                                                     0.06 .
                                    1.1399
                         -1.6578
                                            -1.45
                                                     0.18
             x1
                         -0.0366
                                    0.1004
                                            -0.36
                                                     0.72
             x2
             Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
             Residual standard error: 2.28 on 9 degrees of freedom
             Multiple R-squared: 0.201,
                                         Adjusted R-squared: 0.0231
             F-statistic: 1.13 on 2 and 9 DF, p-value: 0.365
```

(a) Let the dependent variable be the logarithm of Survival time. Fit a multiple linear regression on the independent variables of Mismatch score and Age.

Solution: Based on the program, we can see the  $\beta_0$  equals 11.5724. The estimate of Mismatch score (x1) equals -1.6578. The estimate of Age (x2) equals -0.0366.

(b) Compute an estimate of the variance of the error term.

**Solution:** The residual standard error term will be 2.277 (it is different from the result in screenshot because after several times of running program, the answer is 2.277, not 2.28).

## **Problem 5:**

```
AngerScore, SecondHeartAttack
80, Yes
77, Yes
70, No
68, Yes
64, No
60, Yes
50, Yes
46, No
40, Yes
35, No
30, No
25, Yes
```

Below is the R program used for this problem.

```
# read csv file
mydata <- read.csv("/Users/macbookpro/Desktop/Ex5.csv")</pre>
y <- mydata$SecondHeartAttack;</pre>
x <- mydata$AngerScore;</pre>
# change "YES" or "NO" into booleans
for (i in mydata$SecondHeartAttack){
   if (i == "Yes"){
       i <- 1
   }
   else{
       i <- 0
   }
}
# perform logistic regression on dataset
mylogit \leftarrow glm(y \sim x, data = mydata, family = "binomial")
# show the summary
summary(mylogit)
# predict the new value
newdata <- data.frame(x = 55)
pred <- predict.glm(mylogit, newdata, type = "response")</pre>
pred
```

```
Call:
glm(formula = y \sim x, family = "binomial", data = mydata)
Deviance Residuals:
   Min
            1Q Median
                               30
                                      Max
-1.5201 -1.1511
                 0.7920 0.9932
                                    1.3494
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.04733
                     1.88556 -0.555
                                         0.579
                                         0.444
            0.02606
                     0.03406 0.765
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 16.301 on 11 degrees of freedom
Residual deviance: 15.690 on 10 degrees of freedom
AIC: 19.69
Number of Fisher Scoring iterations: 4
>
```

(a) Explain how the relationship between a second heart attack and one's anger score can be analyzed via a logistic regression model.

**Solution:** Based on the dataset, the values of "Second Heart Attack" contain 'Yes' and 'No'. They can be categorized as two categories. So to some extent, it is a 'Binary Classification' problem. So it can be solved via logistic regression.

- (b) Using a software package of your choice, estimate parameters for this model (for example, in Matlab to fit a logistic model consider the command 'glmfit').
- (c) Estimate the probability that a heart attack patient with an anger score of 55 will have a second heart attack within 5 years.

**Solution:** Based on the program, the probability is 0.5953763.

## **Problem 6:**

(a) For this data set compute the SVD (singular value decomposition) of the original matrix, and using this SVD discuss the expected results of performing a PCA on this data.

Basically the SVD on scaled matrix is the same as PCA. So the SVD on original matrix, I think it is helpless for the PCA.

- (b) Compute the PCA: First compute the mean(s) for the data, and subtract from the original data; second compute the covariance matrix including the scaling 1/(n-1); third compute an eigenvalue decomposition and sort both the eigenvalues and eigenvectors in descending order.
- (c) Plot and discuss the principal components. Discuss how this process and results might differ from a direct SVD of the de-biased, scaled data.

**Solution:** The covariance matrix C is given by  $C = X^T X/(n-1)$ . It is a symmetric matrix so it can be diagonalized as:  $C = V L V^T$ , where V is a matrix of eigenvectors (each column is an eigenvector) and L is a diagonal matrix with eigenvalues *lamda* in the decreasing order on the diagonal.

With Singular Decomposition of X, we can get  $\mathbf{X}=\mathbf{U}\mathbf{S}\mathbf{V}^{T}$ , where S is the diagonal matrix of singular values  $s_{i}$ . So we can use some matrix multiplication rules to get

$$\mathbf{C} = \mathbf{V}\mathbf{S}\mathbf{U}^{\top}\mathbf{U}\mathbf{S}\mathbf{V}^{\top}/(n-1) = \mathbf{V}\frac{\mathbf{S}^2}{n-1}\mathbf{V}^{\top},$$

With the eigenvalue decomposition,  $C=VLV^T$ , so we can obtain  $lamda = s^2/(n-1)$ . So we can know why this process might differ from SVD.

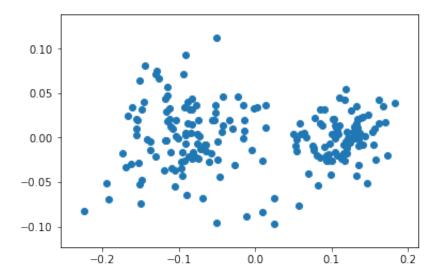
```
In [6]: # read data and import packages
         import math
         import numpy as np
         from scipy import linalg
         from mpl toolkits.mplot3d import Axes3D
         import matplotlib.pyplot as plt
         raw_matrix = numpy.loadtxt(open("/Users/macbookpro/Desktop/IE 529 HW3/Dat
 In [5]: #perform SVD on original matrix
         U,S,V = linalg.svd(raw matrix)
In [13]: #perform matrix scaling and center the data in oringinal matrix
         n = 200
         centered matrix = (raw matrix - raw_matrix.mean(axis=1)[:, None])/ math.s
         cov = np.dot(centered_matrix, centered_matrix.T)
         #calculate the eigenvalue and eigenvectors
         e, v = np.linalg.eig(cov)
         #sort the eigenvalues
         sorted_e = sorted(e , reverse = True)
         vT = v[:,[0,2]]
         print (sorted_e)
         print (e)
         print (v)
         print(vT)
         [[ 0.66943821  0.22771139]
          [ 0.32203254 -0.9467286 ]
          [ 0.66943821  0.22771139]]
         [2.3829744914056272, 0.23466776481580504, 7.1043968626113282e-17]
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             3.22032538e-01
                              5.40520479e-16 -9.46728601e-01]
          ſ
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                                               2.27711392e-01]]
In [11]: renew data= np.dot(vT.T, centered matrix)
         print(renew_data)
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```

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                                    -1.03373071e-02
                                                      -1.82843645e-02
-8.32310220e-03
                   5.43585542e-03
                                    -7.39016057e-03
                                                       1.41140578e-02
 2.09498648e-02
                   1.06737982e-03
                                     1.30013832e-02
                                                      -9.69302791e-03
-1.98490857e-02
                 -2.33831657e-02
                                     7.06782345e-03
                                                       6.35352954e-03
```

```
4.00901696e-03
                     9.61475099e-03
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                                                        2.58120801e-02
   -2.28746541e-03
                    -1.64409843e-02
                                     -1.41867996e-02
                                                       -3.13073564e-02
    2.22028192e-02
                     1.40231167e-02
                                       3.20164616e-02
                                                       -1.92985039e-02
   -2.60580034e-02
                    -1.14489364e-03
                                     -1.78171933e-02
                                                       -1.23310457e-03
   -4.44953945e-03
                     5.66504327e-03
                                     -5.78060561e-03
                                                        1.75413141e-02
   -5.10778719e-02
                     4.29837345e-02
                                       3.23511229e-02
                                                        3.00857802e-02
    2.15105247e-02
                     1.65616145e-02
                                       1.11614030e-02
                                                       -1.47501190e-02
   -2.17320083e-02
                    -1.97758248e-02
                                      -9.21430786e-03
                                                       -1.95444140e-02
   -2.57679258e-02
                     3.57249101e-02
                                       5.41951600e-02
                                                       -2.63268425e-02
    4.28259711e-02
                    -6.60628519e-03
                                       1.59938722e-02
                                                       -4.31412599e-02
   3.32379920e-02
                    -1.44406376e-02
                                     -9.03389486e-03
                                                        1.61501097e-02
   1.16928084e-02
                    -8.32782945e-02
                                     -4.34545987e-02
                                                       -7.67507225e-02
   -3.10188287e-02
                    -9.57984175e-02
                                       4.86936316e-05
                                                       -8.82797396e-02
                    -1.68311804e-03
   -6.82243994e-02
                                     -1.23328459e-03
                                                       -5.39826913e-02
   -9.73705266e-02
                     4.72382976e-03
                                     -4.00754973e-02
                                                       -2.57557909e-02]
]
```

```
In [15]: # plot 1
    X1 = renew_data[0]
    X2 = renew_data[1]

plt.scatter(X1, X2)
    plt.show()
```

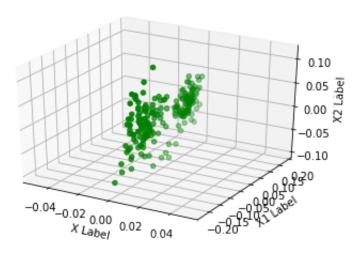


```
In [21]: #plot 2
  plot = plt.figure()
  ax = plot.add_subplot(111, projection='3d')

x = np.zeros(200)
y = X1
z = X2

ax.scatter(x, y, z, c='g', marker='o', label = 'PCA dibased', )

ax.set_xlabel('X Label')
ax.set_ylabel(' X1 Label')
ax.set_zlabel('X2 Label')
plt.show()
```



```
In [27]: # perform SVD on scaled matrix

reconstruct_matrix = centered_matrix
U1, S1, V1 = linalg.svd(reconstruct_matrix)
Ans = np.dot(U1.T, reconstruct_matrix)
print (Ans)
```

```
1.48883676e-01
                     1.01591685e-01
                                       4.89669133e-02
                                                         1.53447482e-01
] ]
    1.92176620e-01
                     1.51254286e-01
                                       8.55679611e-02
                                                         7.24621513e-02
    8.33180676e-02
                     1.15239006e-01
                                       7.47009822e-02
                                                         1.16565833e-01
    5.36900250e-02
                     1.48188761e-01
                                       6.55905856e-02
                                                         1.30247568e-01
    9.55914016e-02
                     5.71234454e-02
                                       4.19264761e-02
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    4.15845085e-02
                     4.90316846e-02
                                       1.15587637e-01
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                                       8.30413808e-02
                                                         9.15261532e-02
    5.55801259e-02
                     4.98841910e-02
                                       6.51740725e-02
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    9.07731440e-02
                     8.43626303e-02
                                       1.53663125e-01
                                                         4.42292880e-02
    7.16734325e-02
                     7.77625621e-02
                                       7.74096271e-02
                                                         7.64352488e-02
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                                       9.47806040e-02
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                     1.37207201e-01
                                       1.05426169e-01
    1.55451769e-01
                     1.17595155e-01
                                       1.11342905e-01
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    9.63610580e-02
                                       2.32566078e-02
                                                         3.38176094e-02
                     1.56057016e-01
    1.08197555e-01
                     1.61999919e-01
                                       1.10503439e-01
                                                         1.11984418e-01
    8.28903941e-02
                     1.30698983e-01
                                       5.98541574e-02
                                                         8.25572392e-02
    1.67061647e-01
                     1.41853711e-01
                                       1.45445316e-01
                                                         1.18306364e-01
    1.15582203e-01
                     6.85233970e-02
                                       2.24726093e-01
                                                         1.44157280e-01
    2.99729943e-02
                     1.60548604e-01
                                       5.25671666e-02
                                                         7.54647646e-02
                                                          02470020- 01
```

```
In [ ]:
```